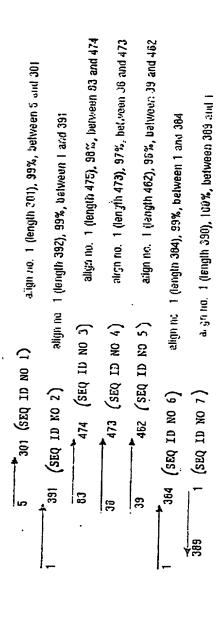
SEQ ID NO 9



FIG

BLASTP ALIGNMENT OF SEQ ID NO: 10, STEM CELL GROWTH FACTOR-LIKE POLYPEPTIDE WITH MOUSE ID NO: THROMBOSPONDIN TYPE 1 DOMAIN POLYPEPTIDE SEQ

Query: Stem ceil growth factor-like plypeptide (SEQ ID NO: 13) Subjet: dbj|BAA75640.1| (AB016768) thrombospondin type 1 domain (Mus musculus) (SEQ ID NO: 25)

Score = 634 (228.2 bits), Expect = 7.8e-62, P = 7.8e-62 Identities = 112/237 (47%), Positives = 152/237 (64%) 19 IGSQNASRGRRQRRNHPNVSQGCQGGCATCSDYNGCLSCKPRLFFALERIGMKQIGVCLS 78 Query:

19 VGSRGI-KGKRORRISABGSQACAKGCELCSBVNGCLKCSPKLFILLERNDIRQVGVCLP 77 SQ C GC CS+ NGCL C P+LF LER ++Q+GVCL +G+RQRR+

Sbjct:

79 SCPSGYYGTRYPJINKCTKCKAD-CDTCFNKYFCTKCKSGFYLHLGKCLDNCPEGLEANN 137 Query:

78 SCPPGYFDARNPDMNKCIKCKIEHCEACFSHNFCTKCQEALYLHKGRCYPACPEGSTAAN 137 SCP GY+ R PD+NKC KCK + C+ CF+ NPCTKC+ YLH G+C Sbjct:

138 HTMECVSIVHCEVSEMNPWSPCTKKGKTCGFKRGTETRVREIIQHPSAKGNLCPPTNETR 197 Query:

138 STWECGSPAQCEMSEWSPWGPCSKRRLCGFRKGSEERTRRVLHAPGGDHTTCSDTKETR 197 TMBC S CE+SEM+PW PC+KK K CGF++G+B R R ++ Sbjct:

198 KCTVQRKKCGKGERGKKGRERKRKKPNKGESKEAIPDSKSLESSKEIPEQRENKQQQ 254 Query:

KCTV+R C -G++ +KG - +R+ N+ +++ +SK S+ Sbjct:

198 KCTVRRTPCPEGQKRRKGGQGRRZNAMRHPARK---NSKEPRSNS----RRHKGQQQ 247

BLASTP ALIGNMENT OF SEQ ID NO: 10, STEM CELL GROWTH FACTOR-LIKE POLYPEPTIDE WITH HUMAN **5**6 POLYPEPTIDE SEQ ID NO: SECRETED PROTEIN CLONE DA228_6

Query: Stem cell growth factor-like plypeptide (SEQ 10 NG: 10) Subjet: sp|W85607|W85607 Secreted protein clone da228_6 (SEQ ID NO: 26) Length = 292

Score = 1477 (525.0 bits), Expect = 2.1e-151, P = 2.1e-151
Identities = 265/265 (100%), Positives = 265/265 (100%)

9 1 MHLRLISWLFIILNFMEYIGSQNASRGRRQRRWFINVSQGCQGGCATCSDYNGCLSCKPR Query:

1 MHLALISWLFIILNFMRYIGSQNASRGRRQRRMHPNVSQGCQGGCATCSDYNGCLSCKPR MHLRLISWLFIILNFMEYIGSQNASRGRRQRRMHPNVSQGCQGGCATCSDYNGCLSCKPR

61 LFFALERIGMKQIGVCLSSCPSGYYGTRYPDINKCTKCKADCDTCFNKNFCTKCKSGFYL 120

LFFALERIGMKQIGVCLSSCPSGYYGTRYPDINKCTKCKADCDTCFNKNFCTKCKSGFYL

Query:

Sbjct:

LFFALERIGMKQIGVCLSSCPSGYYGTRYPDINKCTKCKADCDTCFNKNFCTKCKSGFYL 120 61 Sbjct:

HLGKCLDNCPEGLEANNHTMECVSIVHCEVSEWNPWSPCTKKGKTCGFKRGTETRVREII 180 HLGKCLDNCPEGLEANNHTMECVSIVHCEVSEMNPWSPCTKKGKTCGFKRGTETRVREII 121 Query:

121 HLGKCLDNCPEGLEANNHTMECVSIVHCEVSEANPWSPCTKKGKTCGPKRGTETRVREII 180 Spjet:

240 181 OHPSAKGNLCPPTNETRKCTVQRKKCQKGERGKKGRERKRKKPNKGESKEAIPJSKSLES QHPSAKGNLCPPTNETRKCTVQRKCQKGERGKKGRERKRKKPNKGESKEAI PDSKSLES Query:

QHPSAKGNLCPPTNETRKCTVQRKCQKGERGKKGRERKRKRPNKGESKZAIPDSKSLES 243 181 Shjet:

Query: 241 SKEIPEQRENKQQQKKRKVQDKQKS 265

SKEIPEGRENKQQÇKKRKVQDKQKS

Sojct: 241 SKEIPEQRENKQQQKKRKVQDKQKS 265

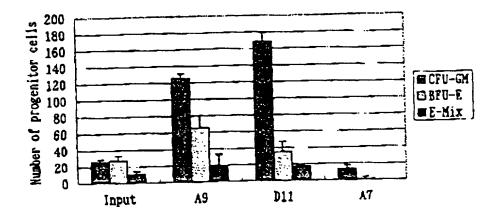


FIGURE 4

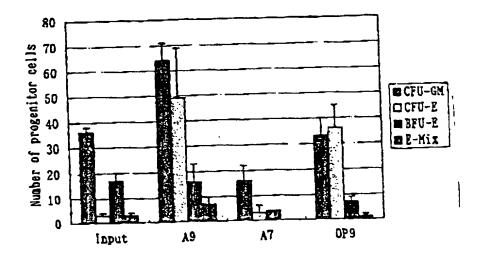


FIGURE 5

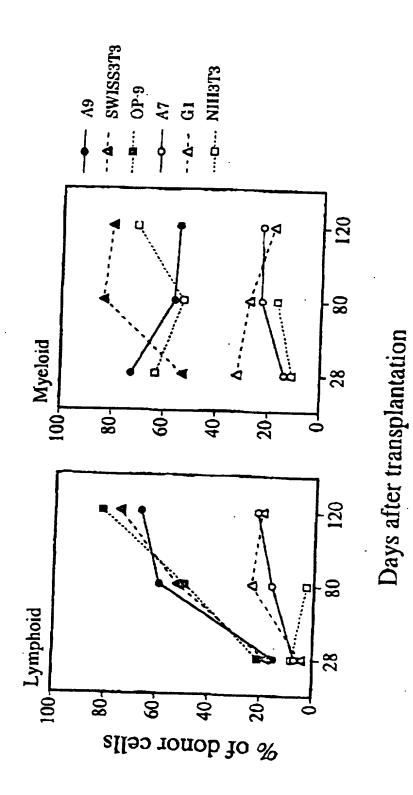
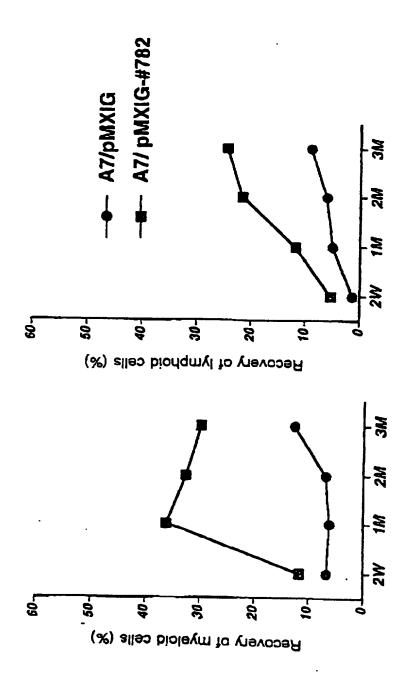


FIGURE 6



Period after transplantation

FIGURE 3